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# Calcium Microdomain Formation at the Perisynaptic Cradle due to NCX Reversal: A Computational Study

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## Abstract

It has recently been proposed using a multi-compartmental mathematical model that negatively fixed charged membrane-associated sites constrain the flow of cations in perisynaptic astroglial processes. This restricted movement of ions between the perisynaptic cradle (PsC), principal astroglial processes and the astrocyte soma gives rise to potassium ( $\text{K}^+$ ) and sodium ( $\text{Na}^+$ ) microdomains at the PsC. The present paper extends the above model to demonstrate that the formation of an  $\text{Na}^+$  microdomain can reverse the  $\text{Na}^+/\text{Ca}^{2+}$  exchanger (NCX) thus providing an additional source of calcium ( $\text{Ca}^{2+}$ ) at the PsC. Results presented clearly show that reversal of the  $\text{Na}^+/\text{Ca}^{2+}$  exchanger is instigated by a glutamate transporter coupled increase in concentration of cytoplasmic  $[\text{Na}^+]_i$  at the PsC, which and instigates  $\text{Ca}^{2+}$  influx through the NCX. As the flow of  $\text{Ca}^{2+}$  along the astrocyte process and away from the PsC is also constrained by  $\text{Ca}^{2+}$  binding proteins, then a  $\text{Ca}^{2+}$  microdomain forms at the PsC. The paper also serves to demonstrate that the EAAT, NKA and NCX represent the minimal requirement necessary and sufficient for the development of a  $\text{Ca}^{2+}$  microdomain and that these mechanisms directly link neuronal activity and glutamate release to the formation of localised  $\text{Na}^+$  and  $\text{Ca}^{2+}$  microdomains signals at the PsC. This local source of  $\text{Ca}^{2+}$  can provide a previously underexplored form of astroglial  $\text{Ca}^{2+}$  signaling.

## 1 Introduction

The concept of astroglial ionic excitability was established in 1990s following the discovery of calcium ion ( $\text{Ca}^{2+}$ ) signaling and propagating  $\text{Ca}^{2+}$  waves in astrocytes in vitro and in situ (Cornell-Bell et al., 1990; Cornell-Bell and Finkbeiner, 1991; Dani et al., 1992; Verkhratsky et al., 1998). Initially, astroglial excitability has been thought to be mediated by  $\text{Ca}^{2+}$  ions; subsequent years have demonstrated signaling roles for sodium ( $\text{Na}^+$ ) (Kirischuk et al., 2012; Parpura and Verkhratsky, 2012; Rose and Chatton, 2016; Rose and Verkhratsky, 2016) and chloride ( $\text{Cl}^-$ ) (Wilson and Mongin, 2018) while a signaling role for potassium ( $\text{K}^+$ ) has also begun to be considered (Breslin et al., 2018), see also (Verkhratsky and Nedergaard, 2018). Ionic signaling in astrocytes serves several physiological roles, in particular, coupling neuronal activity with astroglial homeostatic response within the confines of the astroglial cradle that enwraps at least 50% of all synapses in the central nervous system (Verkhratsky and Nedergaard, 2014; Rose and Verkhratsky, 2016; Verkhratsky and Nedergaard, 2018). Astrocytes express elaborate molecular machinery controlling sodium ( $\text{Na}^+$ ) homeostasis and allowing transient intracellular  $\text{Na}^+$  increases in response to physiological activity in neuronal networks (Kirischuk et al., 2012; Rose and Verkhratsky, 2016). The  $\text{Na}^+$  signals localized to perisynaptic astroglial processes regulate the activity of numerous plasmalemmal transporters responsible for a glutamine-glutamate (GABA) astroglial-neuronal shuttle, for  $\text{K}^+$  buffering, for regulation of pH and for cellular metabolism as well as for secretion of reactive oxygen species scavengers and various neuroactive molecules (Rose and Verkhratsky, 2016; Verkhratsky and Nedergaard, 2016; Verkhratsky and Nedergaard, 2018). Cytoplasmic  $\text{Na}^+$  dynamics are therefore directly linked to the functional activity of astrocytes and represent a mechanism for fast and local signaling at the single synapse/perisynaptic process level. Although cytosolic  $\text{Na}^+$  has emerged as a prominent ion at the interface between signaling and metabolic pathways (Chatton et al., 2016), the spatiotemporal organization of cytosolic  $\text{Na}^+$  dynamics is far from being fully characterized. Additionally, pathways that serve to trigger  $\text{Na}^+$  and  $\text{K}^+$  microdomains need to be resolved.

Computational modelling of astrocyte-neurone interaction is essential for the understanding of the transport processes between cells. However, models of ionic signaling in astrocytes still have some way to go if they are to fully capture the complex morphology of astrocytic processes seen in vivo. Some of these models focus on changes of ionic fluxes crossing the astrocyte membrane controlled by pumps and exchangers, for example  $\text{Na}^+/\text{K}^+$  pump (NKA) and  $\text{Na}^+/\text{Ca}^{2+}$  exchanger, (NCX) and  $\text{K}^+$ ,  $\text{Na}^+$  and  $\text{Cl}^-$  channels as well as axial fluxes of ions in the intracellular space (Haines et al., 2013). This electro-diffusive model allows channels to be distributed in different ways while gating parameters can be altered to match experimental data. Intercellular  $\text{Ca}^{2+}$  waves and oscillations in astrocytes (Pasti et al., 1997; Wade et al., 2011; Wade et al., 2012; Naeem et al., 2015) have also been modelled to identify potential mechanisms of neurone to astrocyte intracellular intercellular signaling.

### 1.1 Astrocytic $\text{Na}^+$ Signaling

The major contributors to astrocyte  $\text{Na}^+$  homeostasis and signaling are the NKA, NCX and the sodium-dependent glutamate transporters (EAAT1 and EAAT2). The NKA regulates resting  $\text{Na}^+$  concentration in the cytosol ( $[\text{Na}^+]_i$ ) and expels excess  $\text{Na}^+$  that enters astrocytes during periods of neuronal activity, whereas glutamate transporters are responsible for the bulk of  $\text{Na}^+$  influx into astrocytes accompanying glutamatergic synaptic transmission (Rose and Verkhratsky, 2016; Rose and Karus, 2013; Kirischuk et al. 2007). The NKA, NCX and glutamate transporters co-localize in

astrocytic perisynaptic processes (Minelli et al., 2007; Melone et al., 2018), suggesting their functional coupling.

All three subunits of NCX (NCX1/SLC8A1, NCX2/SLC8A2 and NCX3/SLC8A3) are expressed in astroglia, with NCX1/SLC8A1 being the predominant isoform (Verkhatsky and Nedergaard, 2018; Pappalardo et al., 2014). The stoichiometry of astroglial NCX is  $3\text{Na}^+:1\text{Ca}^{2+}$ , and hence the equilibrium potential can be calculated from Nernst equation:  $E_{\text{NCX}} = (nE_{\text{Na}} - 2E_{\text{Ca}})/(n - 2)$  where  $n$  is a stoichiometry of  $\text{Na}^+$ , and  $E_{\text{Na}}$  and  $E_{\text{Ca}}$  are equilibrium potentials of  $\text{Na}^+$  and  $\text{Ca}^{2+}$ , respectively. Assuming  $[\text{Ca}^{2+}]_i$  of 50 - 80 nM and  $[\text{Na}^+]_i$  of 15 mM, the  $E_{\text{NCX}}$  could be as negative as  $\sim -85$  to  $-90$  mV, in a similar range to measured resting membrane potential values of astrocytes (Verkhatsky and Nedergaard, 2018). As a result the NCX is prone to fluctuate between forward and reverse transport depending on actual changes in  $[\text{Ca}^{2+}]_i$  and  $[\text{Na}^+]_i$  and the astroglial membrane potential ( $V_m$ ). Conceptually, depolarization or an increase in  $[\text{Na}^+]_i$  will favor NCX operation in the reverse mode, whereas an increase in  $[\text{Ca}^{2+}]_i$  promotes the forward mode of the exchanger. In this way NCX can regulate both  $\text{Ca}^{2+}$  and  $\text{Na}^+$  signals, being relevant in shaping ionic signals in astroglial PsCs.

Recently we presented a new hypothesis (Breslin et al., 2018), which addressed ionic dynamics in thin ( $< 100$  nm) perisynaptic processes whereby negatively charged lipids form deep potential wells near the dipole heads restricting the flow of cations along the process. The negative ion foci serve to form “traps” that attract free cytosolic cations forcing them to hop from trap to trap, thus restricting ion propagation along the process and isolating them from the soma. This “ionic retention by traps” can potentially explain the generation of the transient  $\text{Na}^+$  and  $\text{K}^+$  microdomains at PsCs. The present paper aims to further develop this model with the inclusion of  $\text{Ca}^{2+}$  dynamics at the PsC. We stress here that while this model is not the focus of this paper, it is necessary to include it as it replicates closely the experimentally observed  $\text{Na}^+$  microdomain in astroglial processes (Langer and Rose, 2009). Furthermore, the model allows us to test the hypothesis that reversal of the NCX, due to the  $\text{Na}^+$  microdomain, leads to  $\text{Ca}^{2+}$  influx and this, coupled with  $\text{Ca}^{2+}$  retention along the thin astrocyte process, provides for a plausible mechanism for the generation of a  $\text{Ca}^{2+}$  microdomain at the PsC. Moreover, as this source of  $\text{Ca}^{2+}$  is remote from any endoplasmic reticulum (ER) mediated  $\text{Ca}^{2+}$  release, we propose that this local source of  $\text{Ca}^{2+}$  may provide a previously under-explored form of astrocyte  $\text{Ca}^{2+}$  signaling.

## 2 Model

It was proposed in our earlier paper (Breslin et al., 2018) that  $\text{Na}^+$  and  $\text{K}^+$  retention occurs in thin astrocyte processes. In this paper we consider  $\text{Ca}^{2+}$  retention along the thin processes. This  $\text{Ca}^{2+}$  retention is captured using an extended form of the multi-compartmental mathematical model described in (Breslin et al., 2018). The model consists of a single synapse surrounded by an astrocytic PsC. Due to the complexity of neuronal/astrocytic morphology (Patrushev et al., 2013; Lushnikova et al., 2000; Witcher et al., 2007; Xu-Friedman et al., 2001), this simplified cylindrical compartmental model (Breslin et al., 2018) simplifies the highly complex structures and the associated computational overheads, whilst retaining sufficient functionality to produce meaningful observations. All necessary dimensional details of the model can be found in Table 1. Figure 1 provides details of the multi-compartmental model described by (Breslin et al., 2018).

In the previous study (Breslin et al., 2018),  $\text{K}^+$  and  $\text{Na}^+$  were modelled to determine their influence on ion retention in the thin astrocyte process. In the current study, we adapt this model to include  $\text{Ca}^{2+}$  dynamics within the astrocyte PsC and extracellular space. From Figure 2, it can be seen that the

synapse and PsC contain a number of ionic channels, exchangers and pumps to provide homeostasis and dynamic exchange of ions between the two cells and extracellular space.

The neurone exchanges  $K^+$  and  $Na^+$  with the PsECS via a voltage-gated potassium channel ( $K_{Neu}$ ), voltage-gated sodium channel ( $Na_{Neu}$ ), a sodium potassium pump ( $NKA_{Neu}$ ), a potassium background channel ( $K_B$  on the synapse) and a sodium background channel ( $Na_B$  on the synapse). The astrocyte exchanges  $Na^+$ ,  $K^+$ , and  $Ca^{2+}$  with the PsECS via a background sodium transport ( $Na_B$  on the astrocyte), potassium background transport ( $K_B$  on the astrocyte), potassium inwardly rectifying channel ( $K_{ir}$ ), sodium-potassium-ATPase ( $NKA$ ), a glutamate-sodium-potassium-proton antiporter ( $EAAT1/2$ ), and the  $NCX$ . In both the astrocyte and neurone models, we define  $Na^+$  and  $K^+$  background channels; although modelled as a single transport channel for each ion, these background channels represent a lumped model for  $Na^+$  and  $K^+$  transport, taking into account a multitude of influx and efflux pathways.  $I_{KPF}$ ,  $I_{NaPF}$  and  $I_{CaPF}$  model the flow of  $K^+$ ,  $Na^+$  and  $Ca^{2+}$  ions along the astrocytic process to the soma.  $I_{KECSL}$  and  $I_{NaECSL}$  models the  $K^+$  and  $Na^+$  ionic flow generated by  $K^+$  and  $Na^+$  leaking from the PsECS to the GECS. The mathematical descriptions of ionic exchanges between a neurone and astrocyte at the PsC are now presented. For a more detailed description refer to Breslin et al. (2018).

## 2.1 Astrocyte Model

### 2.1.1 Membrane potential and ion concentrations

In this model the astrocyte resting membrane potential is taken as  $\sim -80$  mV, which is widely reported as the resting membrane potential for astrocytes (Verkhatsky and Nedergaard, 2018). We also assume that isopotentiality is preserved in the PsC as is the case for the astroglial syncytium (Kofuji and Newman, 2004), therefore the astrocyte membrane potential remains fixed at  $\sim -80$  mV.

The perisynaptic model described by (Breslin et al., 2018) comprises five compartments, namely PsC, PsECS, GECS, and the astrocyte process and soma. The astrocyte process is modelled as a long thin cylindrical channel that restricts the flow of cations along the process within the channel due to ion retention. In this work, each of these compartments contains three ionic concentrations,  $K^+$ ,  $Na^+$ , and  $Ca^{2+}$ . All channels, exchangers and transporters permeable to these ions reside on the PsC. The kinetic equations for the changes of ionic concentration of each of these ions is given below. Note:  $z_x F Vol_y$  is used to convert the total ionic current of ion  $x$  into a concentration for the volume  $y$ , where  $z_x$  is the valency of ion  $x$ ,  $F$  is Faradays constant and  $Vol_y$  is the volume of compartment  $y$ . All initial conditions and parameters for this model are described in Tables 2 and 3 respectively. The change in PsC  $K^+$  concentration ( $[K^+]_{PsC}$ ) in the PsC is given by:

$$\frac{d[K^+]_{PsC}}{dt} = - \left( \frac{I_{Kir} + I_{KNKA} + I_{KEAAT} + I_{KPF}}{z_K F Vol_{PsC}} \right)$$

where  $I_{Kir}$  is the  $K_{ir}$  channel current,  $I_{KNKA}$  is the  $K^+$  current through the astrocyte  $NKA$ ,  $I_{KEAAT}$  is the  $K^+$  current created by the glutamate transporter and  $I_{KPF}$  is the  $K^+$  current flowing along the astrocyte process.  $K^+$  changes in the PsECS ( $[K^+]_{PsECS}$ ) is given by:

$$\frac{d[K^+]_{PsECS}}{dt} = - \left( \frac{I_{KECSL} + I_{KNeu} - I_{Km}}{z_K F Vol_{PsECS}} \right)$$

where  $I_{KECSL}$  is current due to  $K^+$  leakage from the PsECS to the GECS,  $I_{KNeu}$  is the  $K^+$  current from the neurone and  $I_{Km}$  is the total  $K^+$  current flowing through the astrocyte membrane.  $K^+$  is held constant at baseline in the GECS and astrocyte soma compartments.

Changes in the PsC  $Na^+$  concentration ( $[Na^+]_{PsC}$ ) is given by:

$$\frac{d[Na^+]_{PsC}}{dt} = - \left( \frac{I_{NaB} + I_{NaNKA} + I_{NaEAAT} + I_{NaNCX} + I_{NaPF}}{z_{Na}FVol_{PsC}} \right)$$

where  $I_{NaB}$  is a current due to  $Na^+$  influx across the membrane via  $Na^+$  permeable ion channels, this is referred to as background  $Na^+$  channel (Breslin et al., 2018),  $I_{NaNKA}$  is the  $Na^+$  dependent current component of the astrocyte NKA,  $I_{NaEAAT}$  is the  $Na^+$  current component of the glutamate transporter,  $I_{NaNCX}$  is the  $Na^+$  current component of the NCX and  $I_{NaPF}$  is the  $Na^+$  current flowing in the astrocyte process.  $[Na^+]$  changes in the PsECS ( $[Na^+]_{PsECS}$ ) is given by:

$$\frac{d[Na^+]_{PsECS}}{dt} = - \left( \frac{I_{NaECSL} + I_{NaNeu} - I_{Nam}}{z_KFVol_{PsECS}} \right)$$

where  $I_{NaECSL}$  is current due to  $Na^+$  leakage from the PsECS to the GECS,  $I_{NaNeu}$  is the  $Na^+$  current from the neurone and  $I_{Nam}$  is the total  $Na^+$  current flowing through the astrocyte membrane.  $Na^+$  is held constant at baseline in the GECS and astrocyte soma compartments.

Changes in the PsC  $Ca^{2+}$  concentration ( $[Ca^{2+}]_{PsC}$ ) is given by:

$$\frac{d[Ca^{2+}]_{PsC}}{dt} = - \left( \frac{I_{CaNCX} + I_{CaPF}}{z_{Ca}FVol_{PsC}} \right)$$

where  $I_{CaNCX}$  is the  $Ca^{2+}$  dependent current component of the NCX, and  $I_{CaPF}$  is the  $Ca^{2+}$  current flowing in the astrocyte process.  $[Ca^{2+}]$  changes in all other compartments are not considered and remain constant at baseline.

### 2.1.2 Glutamate Transporter (EAAT1/2)

Glutamate released into the extracellular PsECS in the course of neurotransmission is assumed here to be entirely removed by astrocytic EAAT1/2. A transport cycle involves the co-transport of 3  $Na^+$  and 1  $H^+$  with 1 glutamate and counter-transport of 1  $K^+$  (Grewer et al., 2014; Murphy-Royal et al., 2015). EAAT1/2 proteins are trafficked to the plasma membrane to facilitate the rapid removal (~3ms) of glutamate from the cleft. Glutamate bound to these proteins is then transported to the astrocytic cytosol over a longer period: in this work a complete transport cycle is assumed to be 30 ms (Otis and Kavanaugh, 2000; Zhou and Danbolt, 2013). Existing EAAT mathematical formulations do not capture adequately this rapid binding and slow release function and therefore in this work we adopt a different approach. To model the stoichiometry and cycle rate we assume that initially a release of glutamate instantaneously binds to membrane-bound proteins and thereafter the flux of  $Na^+$  through the EAAT transporter pore follows an exponentially decaying rate given by:

$$\frac{dJ_{NaEAAT}}{dt} = - \frac{J_{NaEAAT}(t)}{\tau} + J_0\delta(t - tsp)$$

where  $J_{NaEAAT}$  is the flux rate of  $Na^+$  through the EAAT1/2,  $J_0$  is the max flux rate through the transporter,  $\delta$  is the Dirac Delta function,  $t$  is time, and  $tsp$  is the previous neuronal spike time. In our

195 model we view the membrane as a capacitor charged with bound glutamate and  $J_{NaEAAT}$  as a  
196 discharging flux.

197 The  $Na^+$  current through the transporter can be calculated by:

$$198 \quad I_{NaEAAT} = \frac{-J_{NaEAAT} z_{Na} F Vol_{PsECS}}{SA_{PsC}}$$

199 where  $J_{NaEAAT}$  is the  $Na^+$  flux through the EAAT co-transporter,  $z_{Na}$  is the valency of  $Na^+$ ,  $F$  is the  
200 Faradays Constant,  $Vol_{PsECS}$  is the volume of the perisynaptic ECS and  $SA_{PsC}$  is the surface area of  
201 the PsC.

202 The associated  $K^+$  current through the transporter is given by:

$$203 \quad I_{KEAAT} = \frac{-I_{NaEAAT}}{3}$$

204 Note that we are not considering glutamate transport to the astrocytic cytosol as we are only  
205 interested in  $Na^+$ ,  $K^+$  and  $Ca^{2+}$  dynamics. Additionally, our model for EAAT1/2 transport would need  
206 further consideration to include the dependency of fluxes on intra and extra cellular ionic  
207 concentrations.

### 208 **2.1.3 Sodium Calcium Exchanger (NCX)**

209 The NCX is a reversible antiporter which uses the electrochemical gradient of  $Na^+$  to exchange 3  $Na^+$   
210 ions for 1  $Ca^{2+}$  ion across the membrane. Depending on the membrane potential and transmembrane  
211  $Na^+$  gradient, the transporter operates either in forward mode ( $Na^+$  is transported into the cell while  
212  $Ca^{2+}$  is extruded) or in the reversed mode (providing for influx of  $Ca^{2+}$  and efflux of  $Na^+$ ) (Jeffs et al.,  
213 2007).

214 The  $Na^+$  current component of the transporter is given by (Gabbiani and Cox, 2010):

$$215 \quad I_{NaNCX} = \left( \bar{I}_{NCX} \left( \frac{[Na^+]_{PsC}}{[Na^+]_{PsECS}} \right)^3 e^{\frac{\gamma FV}{RT}} - \left( \frac{[Ca^{2+}]_{PsC}}{[Ca^{2+}]_{PsECS}} \right) e^{\frac{(\gamma - 1)FV}{RT}} \right) SA_{PsC}$$

216 where  $\bar{I}_{NCX}$  is the NCX exchanger conductance and  $\gamma$  is a partition parameter.

217 The  $Ca^{2+}$  current component is given by:

$$218 \quad I_{CaNCX} = -2 \left( \frac{I_{NaNCX}}{3} \right)$$

### 219 **2.1.4 Leakage from perisynaptic ECS to global ECS**

220 The diffusion of  $K^+$  and  $Na^+$  between the PsECS and the GECS is modelled as a simple  
221 electrochemical gradient controlled channel in which a zero extracellular potential is assumed and is  
222 given by:

$$223 \quad I_{iECSL} = g_{iECS} E_{iECS} SA_{ECSL}$$

where  $i$  is the ion under consideration,  $g_{iECS}$  is the conductance of the channel,  $SA_{ECSL}$  is the surface area between the PsECS and the GECS, and  $E_{iECS}$  is the Nernst like potential of the channel given by:

$$E_{iECS} = \frac{RT}{F} \ln \left( \frac{[i^+]_{PsECS}}{[i^+]_{GECS}} \right)$$

### 2.1.5 Astrocyte Process Ionic Transport Model

Breslin et al. (2018) proposed that ion retention within thin astrocyte processes can give rise to the formation of  $K^+$  and  $Na^+$  microdomains at the PsC. This localisation of astroglial ionic microdomains arises because in thin processes, surface conduction dominates over volume conduction, and because membrane lipids are negatively charged, deep potential wells form near the dipole heads restricting the flow of cations along the process. Therefore, cations must hop from well to well which restricts ion conduction along the membrane. This hopping effectively semi-isolates the PsC from the astrocytic main body allowing the formation of  $K^+$  and  $Na^+$  microdomains at the PsC under different conditions.

Breslin et al. (2018) proposed that the current flow  $I_{iPF}$  (see figure 2) through the thin process, due to ionic hopping can be represented as:

$$I_{iPF} = K_i \frac{V_A - V_m - V_r}{l} \exp \left[ - \frac{Q_i \left( \phi_w - \sqrt{\frac{Q_i(V_A - V_m - V_r)}{l\pi\epsilon}} \right)}{k_B T} \right] CSA_P$$

where  $K$  is a constant which represents mobility and concentration of mobile ions,  $V_m$  is the resting membrane potential of the astrocyte,  $\phi_w$  is the well activation energy or potential barrier to ion flow,  $l$  is the length of the process,  $Q$  is the charge on a single ion taken as the charge on an electron,  $T$  is the absolute temperature,  $CSA_P$  is the cross-sectional area of the process,  $\epsilon$  is the dynamic permittivity and is given by  $\epsilon = \epsilon_0 \epsilon_r$ , where  $\epsilon_0$  is the absolute permittivity and  $\epsilon_r$  is the relative permittivity of the cytoplasm, and  $k_B$  is the Boltzmann constant.

The concentrations of  $K^+$ ,  $Na^+$ , and  $Ca^{2+}$  in the astrocyte soma are held constant but will be continuously changing at the PsC thus establishing a dynamic concentration gradient associated with these cations. Consequently, we formulate a Nernst-like reversal potential for  $Na^+$ ,  $K^+$  and  $Ca^{2+}$  between the astrocyte soma (AS) and the PsC as:

$$V_r = \frac{RT}{F} \ln \left( \frac{[i]_{AS}}{[i]_{PsC}} \right)$$

where  $i$  is the ion under consideration. A more in-depth discussion and assumptions of the full ionic transport mechanism along thin astrocyte processes is given in (Breslin et al., 2018).

## 2.2 Neurone model

The neuronal model utilized in this work consists of the biophysical Hodgkin and Huxley (HH) type model described in (Breslin et al., 2018) with the addition of a voltage-gated  $Na^+$  channel,  $K^+$  background channel and an  $Na^+$  background channel. These background channels reflect implicit influx/efflux pathways, necessary for the system to ensure dynamic equilibrium. All parameter values for the neurone model are described in Table 4. For reasons of simplicity, the internal concentrations



of neurone  $\text{Na}^+$  and  $\text{K}^+$  remain constant. Whilst we recognize that the neuron NKA is driven by internal  $\text{Na}^+$  and astrocyte NKA is driven by external  $\text{K}^+$ , as we are not considering internal neurone  $\text{Na}^+$  concentration change, we apply the same NKA model for each, altering the maximum pump rates accordingly.

### 2.2.1 Voltage-Gated Neuronal Sodium Channel ( $\text{Na}_{\text{Neu}}$ )

The HH model simulates current flow of  $\text{Na}^+$  through a voltage-gated channel, therefore the current flow of  $\text{Na}^+$  from the neurone can be modelled as:

$$I_{\text{NaNeu}} = -g_{\text{NaNeu}}m^3(V_{\text{Neu}} - E_{\text{NaNeu}})SA_{\text{Syn}}$$

where  $g_{\text{NaNeu}}$  is the maximum  $\text{Na}^+$  channel conductance,  $E_{\text{NaNeu}}$  is the reversal potential of the sodium channel,  $V_{\text{Neu}}$  is the membrane voltage of the neurone and  $SA_{\text{Syn}}$  is the surface area of the synapse.

### 2.2.2 Neuronal Background ion channels ( $\text{Na}_B / \text{K}_B$ )

In this model, there are two individual background ion channels for  $\text{Na}^+$  and  $\text{K}^+$ . These channels use the electrochemical gradient between the PsC and ECS, resulting in an influx of  $\text{Na}^+$  and efflux of  $\text{K}^+$  under normal physiological conditions. They were modelled as a simple passive electrochemical gradient dependent channel given by (Steratt et al., 2011):

$$I_{iB\text{Neu}} = g_{iB\text{Neu}}(V_{\text{neu}} - E_i)SA_{\text{Syn}}$$

where  $i$  is the ion under consideration,  $g_{iB\text{Neu}}$  is the channel conductance. Note: the value of  $g_{iB\text{Neu}}$  is chosen in such a way that the total flux of ion  $i = 0$  at steady state.  $V_{\text{neu}}$  is the neurone membrane voltage,  $SA_{\text{Syn}}$  is the surface area of the neuronal synapse, and  $E_i$  is the channel Nernst potential and is given by:

$$E_i = \frac{RT}{F} \ln \left( \frac{[i^+]_{\text{PsECS}}}{[i^+]_{\text{Syn}}} \right)$$

Note that the concentrations of  $\text{K}^+$  and  $\text{Na}^+$  within the neuronal synapse are held at baseline.

The complete astrocyte/neurone model was implemented using Matlab 2015b 64-bit (Windows version) by Mathworks. All simulation results presented in the results section of this paper used the forward Euler method of integration with a fixed time step of  $\Delta t = 10\mu\text{s}$ .

## 3 Results

This section reports the outcomes of a series of simulations that demonstrate the formation of a  $\text{Ca}^{2+}$  microdomain at the PsC. The simulations show that the  $\text{Ca}^{2+}$  microdomain is a direct result of ion retention along the thin astrocyte process. We have shown in a recent paper (Breslin et al., 2018) that ion retention underpins  $\text{Na}^+$  and  $\text{K}^+$  microdomain formation at the PsC during physiological neuronal excitation. In this work we show that the uptake of  $\text{Na}^+$ , via EAAT channels during neuronal stimulus, creates the  $\text{Na}^+$  microdomain in the PsC thereby causing the NCX to reverse with subsequent formation of a  $\text{Ca}^{2+}$  microdomain in the PsC.

### 3.1 NCX reversal under physiological stimulation

To explore the reversal of the astrocyte NCX, a series of simulations were carried out with the presynaptic neurone stimulated using an external current to produce firing rates of 10 Hz, 20 Hz and 30 Hz respectively. The neuronal stimulus has a duration of ~1 minute where the first 0.1 minute allows the model to reach a steady state condition and the stimulus ceases after 1 min. This long stimulus period allows investigation of what effect sustained neural activity has on the intracellular/extracellular ionic concentrations. In these simulations PsECS  $\text{Ca}^{2+}$  is held constant at baseline, however  $\text{K}^+$  and  $\text{Na}^+$  are permitted to change via the neurone and astrocyte  $\text{K}^+$  and  $\text{Na}^+$  channels. Each time the neurone spikes, it is assumed that there is 1mM of glutamate released into the PsC and there are enough EAAT1/2 transports which allow the instantaneous binding of all the extra glutamate. Therefore, the extracellular glutamate concentration always remains at baseline concentration. Furthermore, the maximum flux rate of EAAT1/2 ( $J_0$ ) is tuned such that 3 mM of  $\text{Na}^+$  is taken up from the PsECs and 1 mM of  $\text{K}^+$  is released into the PsECS by the EAAT1/2 over a period of 30 ms. Moreover, the astrocyte membrane voltage is held constant at ~ -80 mV, in line with reported resting membrane potential for astrocytes, which incidentally is close to the reversal potential for the NCX. Therefore, during periods of homeostatic rest there is no net flow of  $\text{Ca}^{2+}$  or  $\text{Na}^+$  across the membrane associated with the NCX.

The results presented in Figure. 3A show that during periods of neural stimulus,  $\text{K}^+$  ions released by the neurone, build up in the PsECS and are cleared by the astrocyte, which results in a  $\text{K}^+$  microdomain formation at the PsC (Figure 3B). At the start of the neuronal stimulation, there is a transient loss of  $\text{Na}^+$  from the PsECS (Figure 3C), due to neuronal depolarization and astrocytic EAAT transport. Furthermore, the transport of  $\text{Na}^+$  across the astrocyte membrane via  $\text{Na}^+$  background channels and EAAT transporters results in a  $\text{Na}^+$  microdomain formation (Figure 3D).

In addition to  $\text{Na}^+$  and  $\text{K}^+$  microdomain formation, it is clear from Figure 3E that a local PsC  $\text{Ca}^{2+}$  microdomain is also formed. This  $\text{Ca}^{2+}$  microdomain is formed even in the absence of an ER: the ER is widely believed to be essential for astrocyte  $\text{Ca}^{2+}$  dynamics (Verkhratsky et al., 2012). The microdomain of  $\text{Ca}^{2+}$  is caused by the reversal of the NCX causing  $\text{Ca}^{2+}$  influx in exchange for astrocytic  $\text{Na}^+$  efflux. This can be seen in Figure 4C where the only  $\text{Ca}^{2+}$  influx pathway in our model is via the NCX, the complete reversal of the NCX is due to the sudden changes in PsC  $\text{Na}^+$  during neuronal stimulation onset as the astrocyte membrane voltage is held constant.

Figure 4 presents the  $\text{Na}^+$  currents associated with the astrocyte. The main pathway for  $\text{Na}^+$  efflux from the astrocyte is the NKA (Figure. 4 A) while the main pathway responsible for  $\text{Na}^+$  uptake is the EAAT (Figure 4B).

The transient loss of  $\text{Na}^+$ , observed in the PsECS (Figure 3C), is caused by the large, transient uptake by EAAT1/2 accompanying glutamate removal following neuronal stimulation, and the EAAT remains active for approximately 30ms (Figure 4B). This large, transient, EAAT-derived  $\text{Na}^+$  flux, compared to the smaller, slower NKA and NaB activating transport rates, tips the transport uptake/release balance in favor of inward transportation and initiates the formation of a  $\text{Na}^+$  microdomain. As the NKA and NaB uptake increases, due to the increased  $\text{Na}^+$  concentration in the PsC, the influx/efflux,  $\text{Na}^+$  pathways tend towards a state of equilibrium and the microdomain of  $\text{Na}^+$  remains at a stable concentration. It is also noted that as  $\text{Na}^+$  increases in the PsC that the NCX works in reverse mode to remove  $\text{Na}^+$  from the PsC. Figure 4E and F show the  $\text{Na}^+$  currents along the process and between PsECS and GECS respectively. Since the efflux of  $\text{Na}^+$  via the process is several orders of magnitude smaller than the other  $\text{Na}^+$  currents this is the main driving force for the creation of the  $\text{Na}^+$  microdomain in the PsC. Moreover, as the PsECS  $\text{Na}^+$  concentration changes, the

leak current ( $I_{NaECSL}$ ) between the PsECS and GECS attempts to maintain the  $Na^+$  levels in the PsECS.

Figure 5 describes the astrocytic  $Ca^{2+}$  currents; again, it can be seen that during neural stimulation, the influx of  $Na^+$  cause the NCX to work in reverse mode and therefore there is an influx of  $Ca^{2+}$  (Figure 5A). The only efflux pathway for  $Ca^{2+}$  considered within the model is via the thin astrocyte process, which is also governed by the well-hopping mechanism described in (Breslin et al., 2018) (Figure 5B), Therefore the efflux pathway is much less dominant than the influx pathway which results in a microdomain of  $Ca^{2+}$  forming, as seen in Figure 3E.

The  $K^+$  currents behave in a similar manner as previously described in Breslin et al. 2018. NKA and Kir represent the dominant fluxes of  $K^+$  across the astrocytic membrane; NKA is purely responsible for  $K^+$  influx whereas Kir is in a constant state of transition between forward ( $K^+$  influx) and reverse ( $K^+$  efflux) mode during neuronal stimulus (see Figure 6). When neural stimulus ceases, the voltage-dependent reversal of Kir, along with efflux of  $K^+$  via the  $K^+$  background channel, brings the PsC levels of  $K^+$  back to the initial resting state.

#### 4 Discussion

Intracellular ionic signaling represents the substrate for glial excitability (Verkhratsky and Nedergaard, 2018). These intracellular signaling events are mediated through spatially and temporally organized fluctuations in the concentration of major ions; there is firm evidence for physiologically relevant  $Ca^{2+}$  and  $Na^+$  signaling (Rose and Verkhratsky, 2016, Verkhratsky and Nedergaard, 2018; Verkhratsky et al., 2019),  $Cl^-$  signaling (Wilson and Mongin, 2018; Verkhratsky et al., 2019) and  $K^+$  signaling (Olsen et al., 2015; Rimmele and Chatton, 2014). The formation of localized concentration microdomains is critical for ionic signaling spatial fidelity; while the mechanisms underlying formation of these microdomains remain under debate, a new hypothesis has already been proposed (Breslin et al., 2018). In the CNS, most excitatory synapses are tightly enwrapped by perisynaptic astroglial processes forming the synaptic cradle (Reichenbach et al., 2010; Verkhratsky and Nedergaard, 2014). This structure provides homeostatic control of the synaptic cleft and therefore requires the ability to generate relevant signals in response to neuronal activity. The perisynaptic astroglial compartment is devoid of the ER (Reichenbach et al., 2010), hence excluding the metabotropic pathway for generation of local  $Ca^{2+}$  microdomains. Astrocytes express glutamatergic ionotropic receptors, although the receptor-mediated current density is rather low (with whole cell currents rarely exceeding 10 – 100 pA), thus limiting the ionic influx (Verkhratsky and Burnstock, 2014; Rusakov, 2015). Astroglial perisynaptic membranes also express high densities of glutamate transporters (EAAT1/2), which couple glutamate transport with substantial  $Na^+$  influx (Kirischuk et al., 2007; Langer and Rose, 2009). These transporters are co-localized with NCX (Minelli et al., 2007), which couple  $Na^+$  and  $Ca^{2+}$  fluxes in opposite directions. Here we applied the reduced model of PsC to test the hypothesis that glutamate transporters and NCX working together are sufficient to create local  $Ca^{2+}$  microdomains in astroglial perisynaptic cradles.

Our model demonstrates that stimulation of astrocytes with glutamate, mimicking neuronal activity, generates substantial  $Na^+$  influx, which forms local microdomains due to the previously suggested mechanism of ion retention, where cation retention in wells dominates over conventional electrochemical diffusion (Breslin et al., 2018; Wade et al. 2018). Moreover, the generation of  $Ca^{2+}$  microdomains has previously been reported, albeit not at the fine process level, but the underlying ionic fluxes (and channels/transporters contributing to) have not been examined (Rusakov, 2015).

379 This computational modelling study has identified the molecular targets and their relative  
380 contributions to the formation of a  $\text{Ca}^{2+}$  microdomain in the absence of an ER region. Specifically,  
381 our model predicts that the generation of a  $\text{Na}^{+}$  microdomain switches the NCX into reverse mode,  
382 which is sufficient to produce relevant focal  $\text{Ca}^{2+}$  signals; while cross-disciplinary research to test  
383 this hypothesis is beyond the scope of this paper, we envisage that this work will instigate such a  
384 project.

385 In essence only the EAAT, NKA and NCX are necessary and sufficient for the development of a  
386  $\text{Ca}^{2+}$  microdomain: note that to avoid rapidly increasing  $\text{Ca}^{2+}$  concentrations within the cradle, with  
387 increasing neuronal frequency, our model would require a  $\text{Ca}^{2+}$  efflux pathway, and a likely  
388 candidate is the PMCA pump. These mechanisms, which do not depend on intracellular sources for  
389  $\text{Ca}^{2+}$ , directly link neuronal activity and glutamate release to the formation of  $\text{Na}^{+}$  and  $\text{Ca}^{2+}$   
390 microdomains in the perisynaptic astroglial processes, instrumental for generation of astroglial  
391 homeostatic response, which is critical for maintenance of synaptic transmission.

## 392 **5 Conflict of Interest**

393 *The authors declare that the research was conducted in the absence of any commercial or financial*  
394 *relationships that could be construed as a potential conflict of interest.*

## 395 **6 Author Contributions**

396 JW, LM, AV, MD, SH, HVZ, BF and AB contributed to the conception and design of the study. JW  
397 and KB developed the software for simulations. JW and BF created all graphics and data  
398 visualizations. JW, KB, LM, AV, MD, AB wrote the first draft of the manuscript. All authors  
399 contributed to manuscript revision, read and approved the submitted version.

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## 508 **9 Data Availability Statement**

509 No datasets were generated or analyzed for this study.

## 510 **10 Figure Legends**

511 Figure 1. Model Morphology. The model consists of a single synapse enwrapped by a single astrocyte.  
 512 In total there are six compartments, 1) Global Extracellular Space (GECS), 2) Synapse, 3) Perisynaptic  
 513 Extracellular Space (PsECS), 4) Perisynaptic Cradle, 5) Astrocyte Process, and 6) Astrocyte Soma.  
 514 Each compartment is modelled as a cylindrical structure except the GECS and soma, which are deemed  
 515 dimensionless because ionic concentrations remain constant within these compartments (Breslin et al.,  
 516 2018).

517 Figure 2. Ion transport machinery of the perisynaptic cradle and synapse. The model consists of 11  
 518 ionic transports. The synapse contains 5 ionic transports for  $\text{Na}^+$  and  $\text{K}^+$ ; NKA is the sodium/potassium  
 519 pump which extrudes 3  $\text{Na}^+$  ions for every 2  $\text{K}^+$  ions it imports,  $\text{Na}_{\text{neu}}$  and  $\text{K}_{\text{neu}}$  which are the voltage  
 520 gated sodium and potassium channels of the Hodgkin and Huxley model and  $\text{Na}_B$  and  $\text{K}_B$  which are

lump models of all the other  $\text{Na}^+$  and  $\text{K}^+$  channels. The astrocyte contains 6 channels on the inside surface area (facing the synapse) of the perisynaptic cradle; Kir is an inward rectifying  $\text{K}^+$  channel, NKA is a sodium potassium pump similar to that found on the synapse, EAAT1/2 represents the EAAT glutamate transport which is sensitive to glutamate in the synaptic cleft. This transport cotransports 3  $\text{Na}^+$  ions with every glutamate ion and counter transports 1  $\text{K}^+$  ion. Since this work does not consider changes in synaptic and perisynaptic glutamate changes the glutamate ion is not shown in the figure. Moreover, the EAAT dependency on  $\text{H}^+$  is also ignored. The NCX represents the sodium/calcium exchanger which exchanges 3  $\text{Na}^+$  ions for every 1  $\text{Ca}^{2+}$  ion across the membrane. At resting conditions there is a very small exchange in the forward direction as noted in the diagram. This can be reversed under physiological increase of perisynaptic  $\text{Na}^+$  concentration. Finally, the  $\text{Na}_B$  and  $\text{K}_B$  are lumped models of all other  $\text{Na}^+$  and  $\text{K}^+$  ionic transports respectively. The currents  $I_{\text{KECSL}}$  and  $I_{\text{NaECSL}}$  represent leak currents of  $\text{K}^+$  and  $\text{Na}^+$  respectively from the Perisynaptic Extracellular Space (ECS) to the global ECS. It is worth nothing that the currents  $I_{\text{KPF}}$ ,  $I_{\text{NaPF}}$  and  $I_{\text{CaPF}}$  represent the currents of the 3 ions under consideration from the perisynaptic cradle along the process. As described in Breslin et al. (2018), these currents model the hypothesized well-hopping mechanism which severely restricts current flow of these ions along thin processes. This results in the trapping of these ions in the perisynaptic cradle as they are imported across the membrane. Since our model assumes a well-mixed compartment, we consider these trapped ions as a microdomain formation in the perisynaptic cradle.

Figure 3. Astrocyte PsC ion concentrations. (A)  $[\text{K}^+]_{\text{PsECS}}$  transient. (B)  $[\text{K}^+]_{\text{PsC}}$  transient. (C)  $[\text{Na}^+]_{\text{PsECS}}$  transient. (D)  $[\text{Na}^+]_{\text{PsC}}$  transient. (E)  $[\text{Ca}^{2+}]_{\text{PsC}}$  transient. During periods of neuronal activity, increased  $[\text{K}^+]_{\text{PsECS}}$  is cleared by the astrocyte leading to a microdomain of  $\text{K}^+$  in the PsC. Moreover, due to the influx of  $\text{Na}^+$  predominantly via EAAT channels there is an increase in  $[\text{Na}^+]_{\text{PsC}}$ , resulting in the reversal of the NCX and  $\text{Ca}^{2+}$  microdomain formation. The inserts within (A-E) show the fast dynamics of the ionic concentration changes within the PsC in response to the neurone activity.

Figure 4. Astrocyte  $\text{Na}^+$  currents. (A)  $\text{Na}^+$  NKA current. This current is the main pathway for  $\text{Na}^+$  efflux from the astrocyte is the NKA while the main pathway for sodium efflux is  $\text{Na}^+$  current through the EAAT (see (B)) (C)  $\text{Na}^+$  NCX current. It can be seen that during periods of neurone stimulation, the NCX reverses which results in the efflux of  $\text{Na}^+$  from the PsC in response to the increased  $\text{Na}^+$  uptake via the EAAT. (D) Background  $\text{Na}^+$  current. This represents the uptake of  $\text{Na}^+$  through a lumped model of all other  $\text{Na}^+$  channel. The current can be seen to slow down during periods of PsC  $\text{Na}^+$  increase during neurone stimulus. (E)  $\text{Na}^+$  current along the process. The current along the process is governed by the well-hopping mechanism described by Breslin et al. (2018) which severely restricts the flow of  $\text{Na}^+$  from the PsC to the soma. This restriction results in a microdomain of  $\text{Na}^+$  forming in the PsC compartment as the  $\text{Na}^+$  cannot simply diffuse along the process. (F)  $\text{Na}^+$  current due to the leakage of  $\text{Na}^+$  from the PsECS to the GECS which is purely diffusive. The inserts within (A-F) show the fast dynamics of the currents in response to neurone activity.

Figure 5. Astrocyte  $\text{Ca}^{2+}$  currents. (A) Astrocyte  $\text{Ca}^{2+}$  NCX current. During periods of neural stimulation an influx of  $\text{Na}^+$  via the EAAT1/2 cotransporter results in the reversal of the NCX and thus there is an influx of  $\text{Ca}^{2+}$  into the PsC. (B)  $\text{Ca}^{2+}$  current along the process. The only efflux pathway within the model is along the process. The current along the process is governed by the well-hopping mechanism described by Breslin et al. (2018) which severely restricts the flow of  $\text{Ca}^{2+}$  from the PsC to the soma. This restriction results in a microdomain of  $\text{Ca}^{2+}$  forming in the PsC compartment. The inserts within (A-B) show the fast dynamics of the currents in response to neurone activity.

Figure 6. Astrocyte  $\text{K}^+$  currents. (A)  $\text{K}^+$  Kir Current. During periods of neurone stimulation the Kir channel will initially uptake  $\text{K}^+$  before releasing it again at a much slower rate (see insert (A)). (B)  $\text{K}^+$



NKA current. The rate of NKA increases with neurone stimulation due to the increase of PsECS  $K^+$ . This current is mainly responsible for the uptake  $K^+$  in the PsC. (C)  $K^+$  current along the process. Since the current along the process is governed by the well hopping mechanism described by Breslin et al. (2018) it severely restricts the flow of  $K^+$  from the PsC to the soma. This restriction results in a microdomain of  $K^+$  forming in the PsC compartment as the  $K^+$  cannot simply diffuse along the process. Furthermore,  $K^+$  is transported across the membrane via background and EAAT  $K^+$  currents (D, E respectively) much more slowly as these currents are mainly driven by transient  $K^+$  fluctuations in the PsECS due to the neurone stimulation; while these currents have a higher peak magnitude, they are much more “spike like in nature” than NKA currents (See inserts B, D, E). (F)  $K^+$  current due to the leakage of  $K^+$  from the PsECS to the GECS which is purely diffusive.

## 11 Tables

**Table 1: Astrocyte Morphology.**

Parameter	Value	Units	Description
<b>Lengths:</b>			
<b>d<sub>IPS</sub></b>	$300 \times 10^{-9}$	m	Perisynaptic internal diameter
<b>d<sub>EPS</sub></b>	$500 \times 10^{-9}$	m	Perisynaptic external diameter
<b>r<sub>IPS</sub></b>	$150 \times 10^{-9}$	m	Perisynaptic internal radius
<b>r<sub>EPS</sub></b>	$250 \times 10^{-9}$	m	Perisynaptic external radius
<b>l<sub>PS</sub></b>	$300 \times 10^{-9}$	m	Perisynaptic length
<b>d<sub>P</sub></b>	$100 \times 10^{-9}$	m	Process diameter
<b>r<sub>P</sub></b>	$50 \times 10^{-9}$	m	Process radius
<b>l<sub>P</sub></b>	$25 \times 10^{-6}$	m	Process length
<b>d<sub>Syn</sub></b>	$270 \times 10^{-9}$	m	Synapse diameter
<b>r<sub>Syn</sub></b>	$135 \times 10^{-9}$	m	Synapse radius
<b>l<sub>Syn</sub></b>	$300 \times 10^{-9}$	m	Synapse length
<b>Areas:</b>			

<b>CSA<sub>Ps</sub></b>	$3.5343 \times 10^{-14}$	m <sup>2</sup>	Perisynaptic cross sectional area
<b>SA<sub>Ps</sub></b>	$1.4137 \times 10^{-13}$	m <sup>2</sup>	Perisynaptic surface area
<b>CSA<sub>P</sub></b>	$7.854 \times 10^{-15}$	m <sup>2</sup>	Process cross sectional area
<b>SA<sub>P</sub></b>	$7.854 \times 10^{-12}$	m <sup>2</sup>	Process surface area
<b>CSA<sub>Syn</sub></b>	$2.8628 \times 10^{-14}$	m <sup>2</sup>	Synapse cross sectional area
<b>SA<sub>Syn</sub></b>	$1.2723 \times 10^{-13}$	m <sup>2</sup>	Synapse surface area
<b>SA<sub>PsECS-GECS</sub></b>	$1.5715 \times 10^{-14}$	m <sup>2</sup>	Surface area between PsECS and GECS
<b>Volumes:</b>			
<b>Vol<sub>Ps</sub></b>	$1.8850 \times 10^{-17}$	L	Perisynaptic volume
<b>Vol<sub>P</sub></b>	$1.9635 \times 10^{-16}$	L	Process volume
<b>Vol<sub>Syn</sub></b>	$8.5883 \times 10^{-16}$	L	Synapse volume
<b>Vol<sub>PsECS</sub></b>	$2.0145 \times 10^{-18}$	L	Perisynaptic ECS volume

578 **Table 2: Astrocyte Model Variables.**

<b>Variable</b>	<b>Initial Value</b>	<b>Units</b>	<b>Description</b>
<b>V<sub>A</sub></b>	-0.0807	V	Astrocyte Membrane potential
<b>[K<sup>+</sup>]<sub>PsC</sub></b>	0.1	M	K <sup>+</sup> concentration in the perisynaptic cradle
<b>[Na<sup>+</sup>]<sub>PsC</sub></b>	0.015	M	Na <sup>+</sup> concentration in the perisynaptic cradle
<b>[K<sup>+</sup>]<sub>PsECS</sub></b>	0.004	M	Perisynaptic extracellular K <sup>+</sup> concentration
<b>[Na<sup>+</sup>]<sub>PsECS</sub></b>	0.135	M	Perisynaptic extracellular Na <sup>+</sup> concentration
<b>[Glu]<sub>ECS</sub></b>	$25 \times 10^{-9}$	M	Perisynaptic extracellular Glutamate concentration

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580 Table 3: Astrocyte Model Parameters.

Parameter	Value	Units	Description
$V_m$	-0.0807	V	Astrocyte resting membrane potential
$\phi_w$	0.267	eV	Well activation energy
$k_B$	$1.38 \times 10^{-23}$	J/K	Boltzmann constant
$R$	8.31	J/mol/K	Gas constant
$T$	310	K	Temperature
$F$	96485	C/mol	Faraday constant
$Q$	$1.6022 \times 10^{-19}$	C	Coulomb
$C_m$	0.01	F/m <sup>2</sup>	Membrane capacitance
$g_{Kir}$	144	S/m <sup>2</sup>	$K_{ir}$ channel conductance
$g_K$	17.9364	S/m <sup>2</sup>	$K^+$ background transport conductance
$g_{Na}$	0.9761	S/m <sup>2</sup>	$Na^+$ background transport conductance
$K_K$	0.018	S/m	$K^+$ Poole-Frenkel channel constant
$K_{Na}$	0.018	S/m	$Na^+$ Poole-Frenkel channel constant
$PNKA_{max}$	$0.1 \times 10^{-5}$	mol/m <sup>2</sup>	Maximum NKA-ATPase Pump Rate
$K_{Nai}$	$10 \times 10^{-3}$	M	$Na^+$ threshold for NKA-ATPase
$K_{KE}$	$1.5 \times 10^{-3}$	M	$K^+$ threshold for NKA-ATPase
$z_K$	1		$K^+$ Valency

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$z_{Na}$	1		$Na^+$ Valency
$z_{Ca}$	2		Calcium Valency
$[H^+]_{PsC}$	$60 \times 10^{-9}$	M	$H^+$ Concentration in the perisynaptic cradle
$[Glu]_{PsC}$	$1.5 \times 10^{-3}$	M	Glutamate Concentration in the perisynaptic cradle
$[K^+]_{AS}$	0.1	M	$K^+$ Concentration in the astrocyte soma
$[Na^+]_{AS}$	0.015	M	$Na^+$ Concentration in the astrocyte soma
$[Ca^{2+}]_{AS}$	$100 \times 10^{-9}$	M	$Ca^{2+}$ Concentration in the astrocyte soma
$[H^+]_{PsECS}$	$40 \times 10^{-9}$	M	Perisynaptic extracellular $H^+$ concentration
$[Ca^{2+}]_{PsECS}$	$1.5 \times 10^{-3}$	M	Perisynaptic extracellular $Ca^{2+}$ concentration
$[K^+]_{GECS}$	0.004	M	Perisynaptic global ECS $K^+$ concentration
$[Na^+]_{GECS}$	0.135	M	Perisynaptic global ECS $Na^+$ concentration
$[Ca^{2+}]_{GECS}$	$1.5 \times 10^{-3}$	M	Perisynaptic global ECS $Ca^{2+}$ concentration
$[Ca^{2+}]_{PsC}$	$100 \times 10^{-9}$	M	$Ca^{2+}$ concentration in the perisynaptic cradle
$[Ca^{2+}]_{PsECS}$	$1.5 \times 10^{-3}$	M	Perisynaptic extracellular $Ca^{2+}$ concentration
$\epsilon_0$	$8.85 \times 10^{-12}$	F/m	Vacuum permittivity
$\epsilon_r$	0.82	F/m	Relative permittivity of brain tissue
$g_{ECS}$	3.3	S/m <sup>2</sup>	Perisynaptic ECS leak conductance
$\alpha_{EAAT}$	0.0032	A/m <sup>2</sup>	Glutamate transport fitting parameter
$\beta_{EAAT}$	28.8	mV <sup>-1</sup>	Glutamate transport fitting parameter

$r_g$	$5 \times 10^{-7}$	$M^{-1}$	Slope of glutamate uptake
$s_g$	$9 \times 10^{-6}$	M	Threshold for glutamate uptake
$\bar{I}_{NCX}$	1	A/m <sup>2</sup>	NCX Partition Parameter
$\gamma$	0.5		
$J_0$	0.06	M/s	Maximum EAAT1/2 Flux rate

581 **Table 4: Neurone Parameters.**

Parameter	Value	Units	Description
$P_{NKAmaxNeu}$	$-3.7863 \times 10^{-8}$	mol/m <sup>2</sup>	Maximum NKA-ATPase Pump Rate
$K_{NaNeu}$	$10 \times 10^{-3}$	M	Na <sup>+</sup> threshold for NKA-ATPase
$K_{KNeu}$	$1.5 \times 10^{-3}$	M	K <sup>+</sup> threshold for NKA-ATPase
$[Na^+]_{Syn}$	0.015	M	Na <sup>+</sup> concentration in the synapse
$[K^+]_{Syn}$	0.1	M	K <sup>+</sup> concentration in the synapse
$g_{KNeu}$	360	S/m <sup>2</sup>	Maximum K <sup>+</sup> channel conductance
$g_{NaNeu}$	1200	S/m <sup>2</sup>	Maximum Na <sup>+</sup> channel conductance
$g_{LNeu}$	3	S/m <sup>2</sup>	Maximum leak channel conductance
$g_{KBNeu}$	1.0522	S/m <sup>2</sup>	K <sup>+</sup> Background channel conductance
$g_{NaBNeu}$	2.3217	S/m <sup>2</sup>	Na <sup>+</sup> Background channel conductance
$E_{KNeu}$	-0.12	V	K <sup>+</sup> channel reversal potential
$E_{NaNeu}$	0.115	V	Na <sup>+</sup> channel reversal potential
$E_{LNeu}$	0.010613	V	Leak channel reversal potential

<b>C<sub>m</sub></b>	0.01	F/m <sup>2</sup>	Membrane capacitance
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